IPHC 5-year Research Program

RAB Meeting November 16, 2016

Josep V. Planas Biological & Ecosystem Science Program Manager

Planned research activities at IPHC



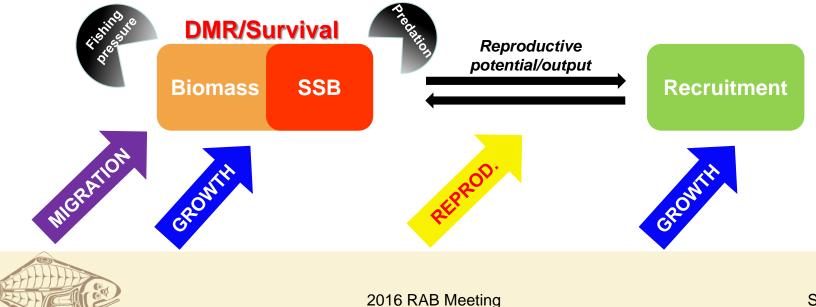


INTERNATIONAL PACIFIC HALIBUT COMMISSION

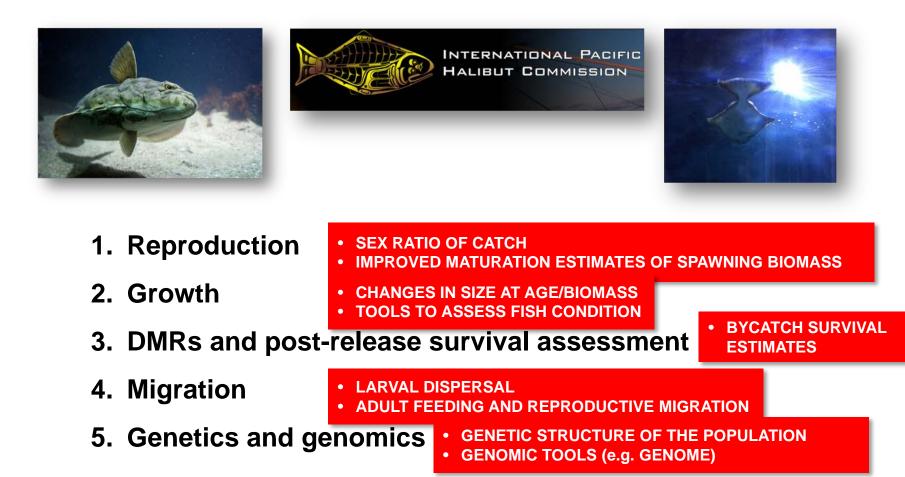


Primary objectives

- Identify and address critical knowledge gaps in the biology of the Pacific halibut
- Understand the influence of environmental conditions on halibut biology
- Apply resulting knowledge to reduce *uncertainty* in current stock assessment models



Planned research activities at IPHC





1. Reproduction: proposed studies

There are important knowledge gaps on the reproductive biology of the species

- SEX RATIO OF CATCH
- IMPROVED MATURATION ESTIMATES OF SPAWNING BIOMASS

What is needed?

- Knowledge on reproductive development, maturation, fecundity, sex determination mechanisms (sex identification), environmental and hormonal control of reproduction.
- Scientific-based criteria to identify reproductive status and potential.
- Updated estimates of age and size at maturation.
- Information on skipped spawning.

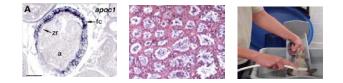
- Full characterization of the annual reproductive cycle
- Identification of sex determination mechanism(s) and influencing factors

1. Reproduction: proposed studies

• Full characterization of the annual reproductive cycle

Objective: Understand temporal changes in reproductive development throughout an entire annual reproductive cycle in male and female Pacific halibut

- Histological assessment of gonadal development and maturation.
- Endocrine profiling (hormone levels in the blood) of the reproductive cycle.
- Gene expression (transcriptome) profiling of the reproductive axis.
- Gonadosomatic index (GSI) measures throughout the reproductive cycle.
- Ultrasound monitoring of gonadal development and maturation.



Deliverables:

- Comprehensive reproductive monitoring of the adult population in order to improve our estimate of actual spawning biomass
- Accurate staging of reproductive status
- Updated maturity-at-age estimates.
- Estimates of skipped-spawning



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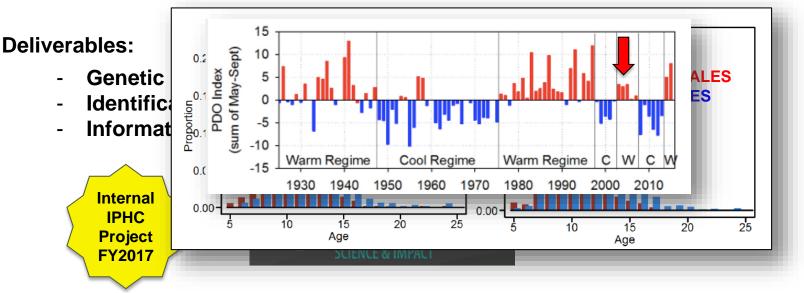
Project FY2017

1. Reproduction: proposed studies

• Identification of sex determination mechanisms

Objective: Understand how sex is established in Pacific halibut

- Identification of sex markers (SNPs)
- Identification of the sex determining mechanism(s) and its onset during early development.
- Identification of environmental influences (e.g. temperature) on sex determination.
- Evaluate possible consequences on sex ratios at the population level.



Little is known regarding what factors influence growth in this species

- CHANGES IN SIZE AT AGE/BIOMASS
- TOOLS TO ASSESS FISH CONDITION

What is needed?

- Knowledge on growth patterns and environmental influences.
- Improved understanding in the possible role of growth alterations in the observed decrease in size at age.

- Extensive catalogue of physiological markers to monitor growth
- Evaluation of growth patterns and effects of environmental influences



• Extensive catalogue of molecular markers for growth

Objective: Identify and validate molecular growth-related markers for growth studies.

- Identification of expressed sequences from skeletal muscle (white and red) and liver.
- Develop molecular assays to quantify gene expression of growth markers in relevant tissues.



3. De novo Transcriptome Assembly Stats

Sample ID	Total trinity 'genes'	Total trinity transcripts	Percent GC	Contig N50	Median contig length	Average contig	Total assembled bases	
R116-pool1	37,161	39,638	47.76	1,198	385	721.49	28,598,382	WHITE MUSCLE
R116-pool2	38,143	40,814	46.02	1,096	398	691.85	28,237,340	LIVER
R116-pool5	70,693	86,561	47.17	2,104	495	1,051.87	91,050,930	RED MUSCLE

5.1 Mapping statistics

Sample ID	Danio rerio	uniprot	est others	total	unmapped	Danio%	uniprot%	est others%	unmapped%	
R116-pool1	13,873	2,661	23,066	39,638	38	35.00%	6.71%	58.19%	0.10%	WHITE MUSCLE
R116-pool2	13,233	2,547	24,998	40,814	36	32.42%	6.24%	61.25%	0.09%	LIVER
R116-pool5	25,341	5,579	55,466	86,561	175	29.28%	6.45%	64.08%	0.20%	RED MUSCLE

Deliverables:

- Establishment of a growth-related gene sequence dataset
- Molecular assays to monitor growth patterns



Evaluation of growth patterns and effects of environmental influences

Objective: Identify molecular, biochemical and isotopic profiles characteristic of specific growth patterns and evaluate potential effects of environmental influences.

- Evaluation of different growth trajectories in the wild.
 - In BS NMFS trawl survey in 2016:
 - 75 fish <40 cm length
 - 75 fish 40-60 cm length
 - 75 fish 60-80 cm length



Characterization of molecular and biochemical growth markers in liver and muscle samples from age-matched individuals

Establishment of different growth trajectories in juvenile fish in captivity to identify molecular and biochemical signatures of growth patterns.



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High rate ormal rate

field Marine Science Cente

Isotopic tissue turnover to trace dietary and/or habitat shifts





- Investigate the effects of environmental factors on growth performance.
 - Effects of temperature, salinity, dissolved oxygen and water pH on growth.
 - Identify the optimal environmental conditions for growth.
- Understand the basis of the sexual dimorphic growth in the Pacific halibut.

Deliverables:

- Identification and validation of growth markers for field studies
- Characterization of molecular and biochemical growth signatures
- Environmental effects on somatic growth
- Improved biological inputs on biomass estimates





Little is known regarding what factors influence bycatch survival and how

• BYCATCH SURVIVAL ESTIMATES

What is needed?

 To introduce quantitative measurable factors that are linked to fish handling practices and to fish physiological condition and ultimately to survival in order to improve current DMR estimations

- Evaluation of the effects of fish handling practices on injury levels and the physiological condition of captured Pacific halibut
- Investigate the relationship between physiological condition post-capture and survival as assessed by the use of accelerometer tags.
- Improving estimates of survival of Pacific halibut caught in the trawl fishery

• Evaluation of the effects of fish handling practices on injury levels and the physiological condition of captured Pacific halibut

Objective: Understand relationship between handling practices and physiological condition of captured Pacific halibut in the longline fishery

- Assess injuries associated with release techniques (gangion cut, careful shake, hook straigthening).
- Determine the physiological condition of all captured fish with associated injury levels after different deck exposure times: condition factor index (Kn), energy (fat) levels, morphometric analyses.
- Measure the levels of stress and physiological disturbance indicators in the blood of all captured fish (cortisol, lactate, glucose, potassium, hematocrit).

Deliverables:

- Injury profile for different release techniques in the longline fishery
- Physiological assessment of fish handling practices: fish condition index post-capture



 Investigate the relationship between physiological condition postcapture and survival as assessed by tagging.

Objective: Measure survival post-release in Pacific halibut and link this with the physiological condition and capture-related events

- Tag fish that have been exposed to different handling practices in the longline fishery with accelerometer tags in addition to conventional tags (wire).
- Assess survival of fish according to size and physiological conditon.

Deliverables:

- Information on post-release survival in relation to handling practices and physiological condition.
- Information on post-release survival in relation to size.

Hook

Injury

Code

- Estimating DMRs by EM.

In

Hook

Release

Technique

Handling events

Hook Release lechnique Survival

• Improving estimates of survival of Pacific halibut in the trawl fishery

Objective: Assess tl non-directed trawl f

- Continue an collaborativ
- Apply method
- Determine s
- Relate physi

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arded Pacific halibut in the of discard mortality rates

Amendment 80 fleet) to plan

ptured halibut.

scarded halibut

Deliverables:

 Improved knowledge of survival of discarded halibut and, consequently, improved estimates of discard mortality rates in the trawl fishery.



4. Migration: proposed studies

- LARVAL DISPERSAL
- ADULT FEEDING AND REPRODUCTIVE MIGRATION

What is needed?

- Improve our understanding on larval, juvenile and reproductive migration.
- Incorporate additional sources of biological information on migratory.

- Towards a more integrative view on migration
- Larval migration and connectivity
- Swimming and migratory performance



4. Migration: proposed studies

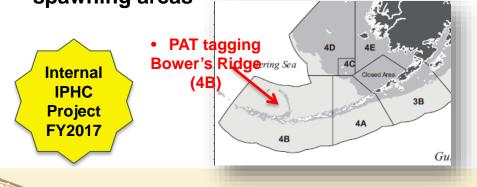
• Towards a more integrative view on migration

Objective: Combine current tagging efforts with genetic and otolith and tissue composition analyses.

- Genetic analyses of tagged fish to shed light on migration patterns and geographic origin.
- Otolith microchemical and stable isotope analyses and tissue stable isotope analyses.
- Reproductive monitoring of PAT-tagged adult females: blood endocrine reproductive parameters, ovarian tissue biopsies and ultrasound for ovarian staging.

Deliverables:

- Genetic and elemental and isotopic information on migratory adult fish
- Improved knowledge on reproductive migrations and identification of spawning areas



• Tail pattern recognition

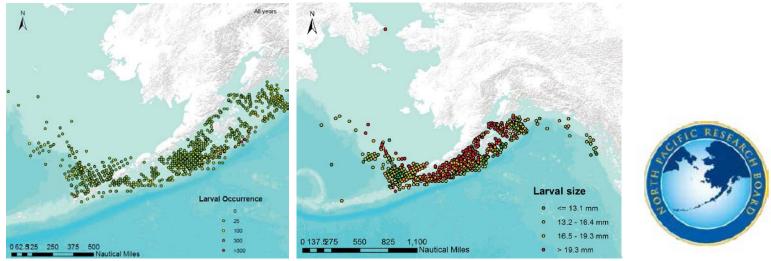


4. Migration: proposed studies

• Larval migration and connectivity

Objective: Understand the mechanisms of larval connectivity between the GOA and the BS.

- Collect data from the NMFS icthyoplankton survey and map larval distribution over time and space.
- Collect larval samples from the survey to conduct genetic analyses.



Deliverables:

Collaboration with Janet Duffy-Anderson, Esther Goldstein, William Stockhausen (NOAA-AFSC)

- Improved knowledge on larval distribution, migration and genetic structure within the population

5. Genetics and genomics: proposed studies

GENETIC STRUCTURE OF THE POPULATION
GENOMIC TOOLS (e.g. GENOME)

What is needed?

- Improved knowledge on the genetic composition of the population
- Establish genomic resources for the species
- Genome-wide association studies to evaluate genetic effects of fisherydependent and fishery-independent influences on growth, reproduction, nutrition, etc.

- Population genetic studies
- Sequencing of the Pacific halibut genome

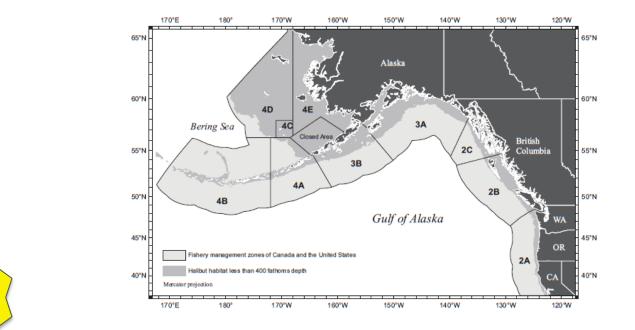


5. Genetics and genomics: proposed studies

Population genetic studies

Objective: Genetic characterization of Pacific halibut throughout its distribution range

- Characterization of population structure by RAD sequencing and SNP analysis.
- Identification of genetic signatures of geographical population groups





Internal

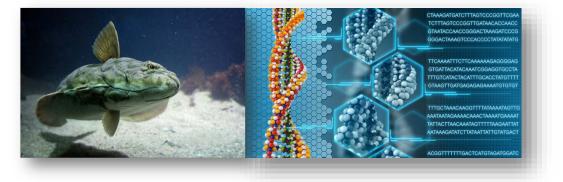
IPHC Project

FY2017

5. Genetics and genomics: proposed studies

• Pacific halibut genome

Objective: Obtain a first draft sequence of the Pacific halibut genome



- Identify genomic regions and genes responsible for temporal and spatial adaptive characteristics.
- Genome-wide association studies to try to understand the genetic basis of growth, reproductive performance, migratory behaviour and performance, etc.
- Provide genomic resolution to genetic markers (from RAD tag seq or RNAseq).

Internal IPHC Project FY2017



Link genotype and phenotype.

Temporal chart of activities

2016	2017	2018	2019	2020	2021				
	Annual repro	ductive cycle							
	Sex determinatio	n mechanisms							
Рори	llation genetic stru	cture							
	Genome s	equencing							
		Genom	e-wide association	studies					
	Epige	enome characteriz	ation						
Growth trai	nscriptome								
Gro	owth-related patter	rns							
	Regi	ulation of growth by	y environmental fa	ctors					
	Handling praction	ces, injury levels							
	and physiolog	gical condition							
		Physiological	condition and						
		surv	/ival						
	Comprehensive studies on migration								
Tagging sublegal halibut									
Reproductive monitoring of PAT-tagged adults									



